

Image motion

## finding a template

Suppose we wish to find a known template T(x,y) in a given image I(x,y).

This problem is known as template matching.



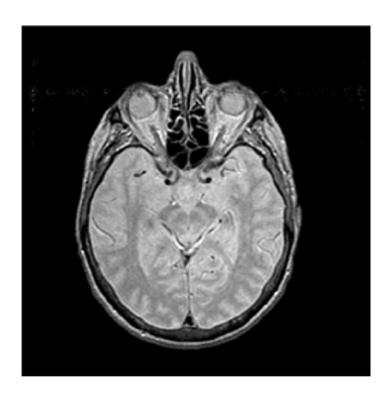
template

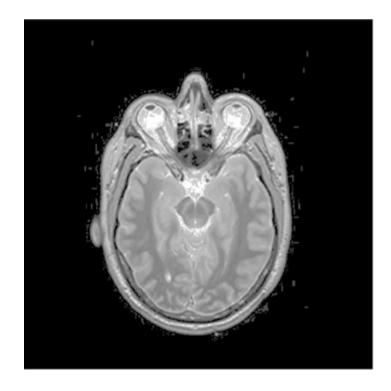
image

**FC Barcelona** 

the template can be small or large

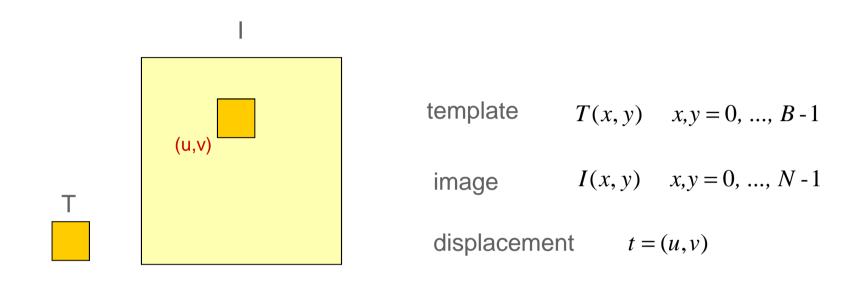
# alignment (MRI images)





atlas test slice

## template matching

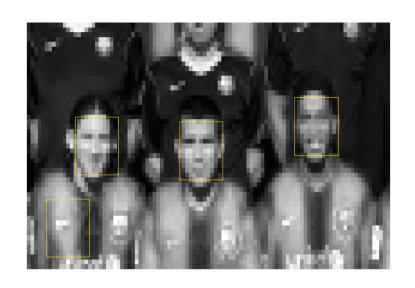


The solution is based on two steps:

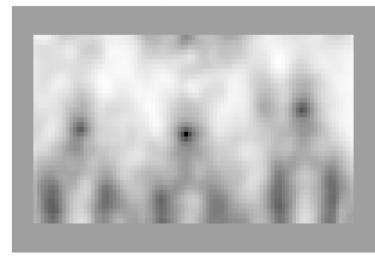
- define a matching criterion M
- find local maxima/minima

- (e.g., cross correlation)
- (e.g., exhaustive search)

## object detection



matching criterion: M



nonlinear optimization



Non-minimum suppression:

 $M(t_0) < M(t)$  for all t in a vicinity of radius r of d

Thresholding:

 $M(t_0) < \lambda$ 

 $\lambda$  threshold

### matching criteria

cross-correlation

$$R(u,v) = \sum_{x,y=0}^{B-1} T(x,y)I(x+u,y+v)$$

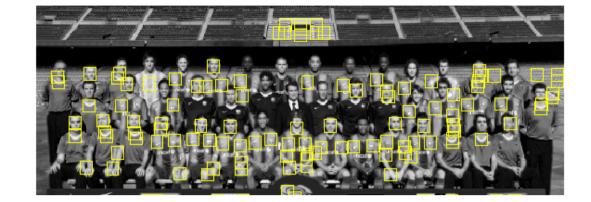
sum of square differences (SSD) (l<sub>2</sub> norm, squared)

$$E(u,v) = \sum_{x,y=0}^{B-1} [T(x,y) - I(x+u,y+v)]^2$$

sum of absolute differences (SAD) (I<sub>1</sub> norm)

$$E(u,v) = \sum_{x,y=0}^{B-1} |T(x,y) - I(x+u,y+v)|$$

Non integer displacements can be considered. Image interpolation is required in this case.



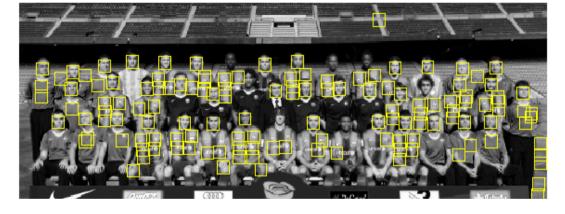
cross-correlation

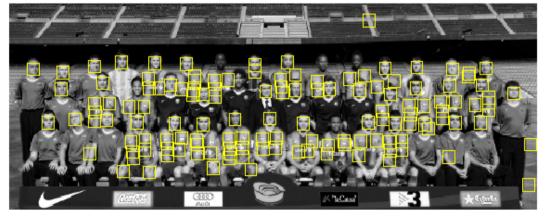
SSD

4

SAD

•





### limitations



X

### Template matching has weaknesses:

- not invariant to rotations and scaling
- not invariant to illumination changes
- time consuming
- template adaptation is tricky

- → more general transformations
- modify matching criteria to improve robustness

### problem formulation





Matlab

### Image alignment

Given 2 (or more) images I, T we wish to estimate a transformation which maps the first into the second

$$(x, y) \rightarrow (x', y')$$
  $(x', y') = W(x, y; \theta)$ 

according to some criterion.

This can be done using:

#### feature based methods:

based on the alignment of feature points (marks)

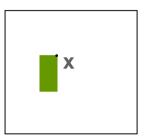
#### image based methods:

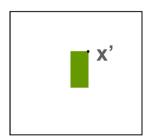
based on the alignment of image intensity or color

What geometric transformations can we use?

## translation & rigid body

#### translation



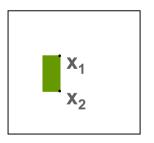


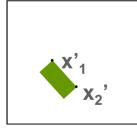
$$W(x;\theta) = x + t$$

$$\theta = t$$

2 degrees of freedom

### rigid body





$$W(x;\theta) = Rx + t$$

 $\theta = (R, t)$ 

3 degrees of freedom

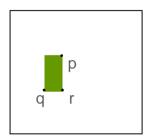
rotation matrix

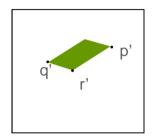
$$RR^{T} = R^{T}R = I$$
  
 $det(R) = 1$ 

$$\mathbf{R} = \begin{bmatrix} \cos \theta & -\sin \theta \\ \sin \theta & \cos \theta \end{bmatrix}$$

### afinne and projective transformations

#### affine transformation

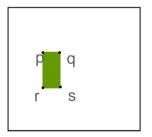


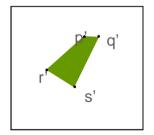


$$W(x;\theta) = Ax + t$$
  $\theta = (A,t)$ 

6 degrees of freedom

#### projective transformation (homography)





$$W(x,\theta) = \begin{bmatrix} \frac{p_1x + p_2y + p_3}{p_7x + p_8y + p_9} \\ \frac{p_4x + p_5y + p_6}{p_7x + p_8y + p_9} \end{bmatrix} \qquad \theta = (p_1,...,p_9)$$

8 degrees of freedom

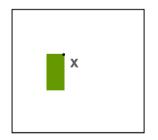
## projective and polynomial transformations

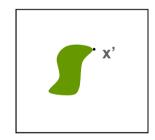
#### projective (contd.)

$$x' = \frac{\widetilde{\mathbf{x}}^{\mathrm{T}} \mathbf{p}_{1}}{\widetilde{\mathbf{x}}^{\mathrm{T}} \mathbf{p}_{3}} \qquad y' = \frac{\widetilde{\mathbf{x}}^{\mathrm{T}} \mathbf{p}_{2}}{\widetilde{\mathbf{x}}^{\mathrm{T}} \mathbf{p}_{3}}$$

$$p_1 = [p_1 p_2 p_3]^T$$
  $p_2 = [p_4 p_5 p_6]^T$   
 $p_3 = [p_7 p_8 p_9]^T$   $\tilde{x} = [x y 1]^T$ 

### polynomial





$$\mathbf{W}(\mathbf{x}, \boldsymbol{\theta}) = \begin{bmatrix} \sum_{p,q: p+q \le n} a_{pq} x^p y^q \\ \sum_{p,q: p+q \le n} b_{pq} x^p y^q \end{bmatrix}$$

The estimation of coefficients is numerically ill conditioned

others .... e.g., free form deformations

# properties

	DoF	Preserves lines?	Preserves Paralelism?	Preserves Angles?	Preserves length?
translation	2	Yes	Yes	Yes	Yes
Rigid body	3	Yes	Yes	Yes	Yes
Affine	6	Yes	Yes	X	X
Projective	8	Yes	X	X	X
Polynomial	(n+2)(n+1)/2	X	X	X	X

can we align images using intensity?

### image based methods

#### Problem:

Given two images T, I we wish to find a geometric transformation W(x) which maps points of the first image into points of the second, such that  $I(W(x)) \approx T(x)$ .

Most popular criterion (SSD)

$$E(\theta) = \sum_{\mathbf{x}} [T(\mathbf{x}) - I(W(\mathbf{x}; \theta))]^{2}$$

Note: the sum is for all the points x in which both images T(x), I(W(x)) overlap.

The minimization of E is a non linear problem!!

## Lucas-Kanade (translation motion)

Criterion 
$$E(u,v) = \sum_{\mathbf{x}} [T(\mathbf{x}) - I(\mathbf{x} + \mathbf{t})]^2$$

Parameter update  $t = t_0 + \Delta t$ 

First order approximation of the image

$$I(\mathbf{x} + \mathbf{t}) = I(\mathbf{x} + \mathbf{t}_0) + \nabla I(\mathbf{x} + \mathbf{t}_0)^T \Delta \mathbf{t}$$

#### Lucas Kanade algorithm (recursion)

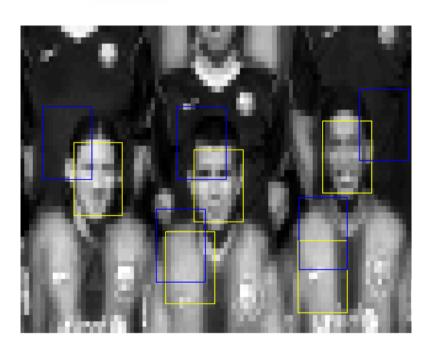
$$\begin{bmatrix} \sum I_x^2 & \sum I_x I_y \\ \sum I_y I_x & I_y^2 \end{bmatrix} \Delta t = \begin{bmatrix} \sum (T(x) - I(x + t_0))I_x \\ \sum (T(x) - I(x + t_0))I_y \end{bmatrix}$$
$$t \leftarrow t_0 + \Delta t$$

$$R\Delta t = r$$
$$t \leftarrow t_0 + \Delta t$$

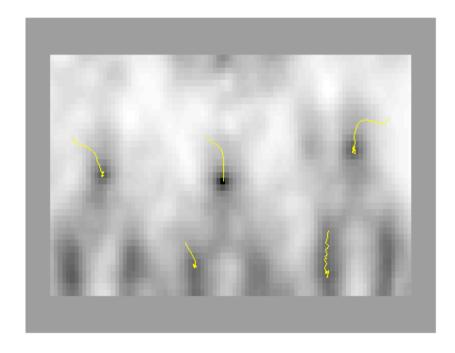
 $I_x$ ,  $I_y$  are the partial derivatives of I at  $x+t_0$ .

# convergence from several starting points





SSD criterion



The SSD criterion is not explicitly computed in the L-K algorithm.

### proof

Let us minimize

$$E = \sum_{x} [T(x) - I(x + t_0) - \nabla I(x + t_0)^T \Delta t]^2$$

A necessary condition is

$$\frac{dE}{d\Delta t} = 0 \qquad \sum_{x} [T(x) - I(x + t_0) - \nabla I(x + t_0)^T \Delta t] \nabla I(x + t_0) = 0$$

$$\sum_{x} \nabla I(x+t_0) \nabla I(x+t_0)^T \Delta t = \sum_{x} [T(x) - I(x+t_0)] \nabla I(x+t_0)$$

Defining

$$\nabla I(\mathbf{x} + \mathbf{t}_0) = \begin{bmatrix} I_x(\mathbf{x} + \mathbf{t}_0) \\ I_y(\mathbf{x} + \mathbf{t}_0) \end{bmatrix}$$

We obtain

$$\begin{bmatrix} \sum I_x^2 & \sum I_x I_y \\ \sum I_y I_x & \sum I_y^2 \end{bmatrix} \Delta t = \begin{bmatrix} \sum (T(x) - I(x + t_0))I_x \\ \sum (T(x) - I(x + t_0))I_y \end{bmatrix}$$

### discussion

#### L-K strong points

- uses all the available information
- It is simple
- appropriate for tracking
- can be extended to deal with general motion models

#### L-K weak points

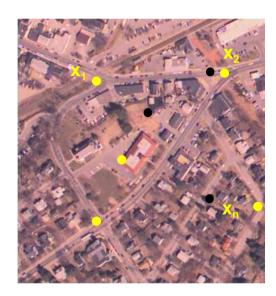
- no guarantee that the optimal solution is obtained
- the solution depends on the initialization —— use multiple scales
- convergence is difficult if the number of parameters is high
- solution depends on the illumination
   Illumination can be estimated

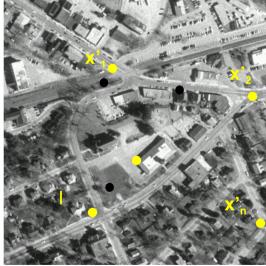
can we align images from sparse prototypes?

### feature based matching

#### Problem:

Given two sets of points  $\{x_i\}$ ,  $\{x_i'\}$  detected in the images T, I, we wish to find a geometric transformation W that maps the points  $\{x_i\}$  into the points  $\{x_i'\}$ .





$$\mathbf{x}_{i} = \begin{bmatrix} x_{i} \\ y_{i} \end{bmatrix}^{T} \qquad \mathbf{x'}_{i} = \begin{bmatrix} x_{i'} \\ y_{i'} \end{bmatrix}$$

we assume that the correspondence is known

## approach

Define a matching criterion e.g.,

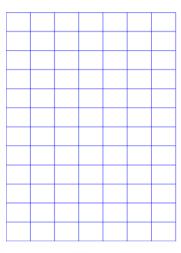
$$E(\theta) = \sum_{i} \|\mathbf{x'_i} - \mathbf{W}(\mathbf{x_i}; \theta)\|^2$$
 SSD criterion

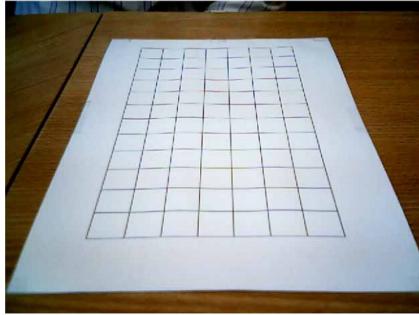
Minimize the criterion with respect to  $\theta$  using a closed form or a numeric algorithm.

Note: there are other matching e.g., I<sub>1</sub>norm.

# example

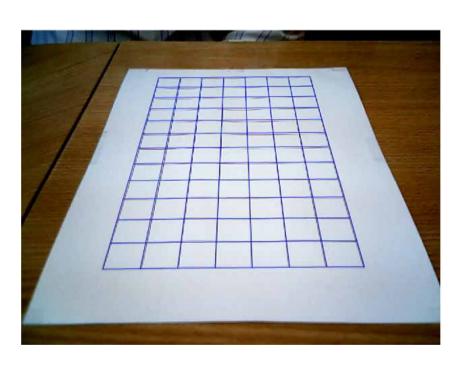
input





Jorge Marques, 2008

#### output



alignment using a projective transform

### estimation of an homography

### Homography

$$x' = f(x, p)$$

$$x' = \frac{p_1 x + p_2 y + p_3}{p_7 x + p_8 y + p_9}$$
$$y' = \frac{p_4 x + p_5 y + p_6}{p_7 x + p_8 y + p_9}$$

||p||=1

is a nonlinear function of the unknown parameters.

The minimization of the SSD criterion is difficult!!

$$E(p) = \sum_{i} \|x'_{i} - f(x_{i}, p)\|^{2}$$

Idea: use another (simpler) criterion instead

$$(p_7x + p_8y + p_9)x' = (p_1x + p_2y + p_3)$$
$$(p_7x + p_8y + p_9)y' = (p_4x + p_5y + p_6)$$

$$e = \begin{bmatrix} (p_1x + p_2y + p_3) - (p_7x + p_8y + p_9)x' \\ (p_4x + p_5y + p_6) - (p_7x + p_8y + p_9)y' \end{bmatrix}$$

algebraic error

$$E'(p) = \sum_{i} \|e_{i}\|^{2} \|p\| = 1$$

## estimation of the projective transform (2)

minimize

$$E' = p^{T}M^{T}Mp$$

with restriction  $p^T p = 1$ 

$$M = \begin{bmatrix} x_1 & y_1 & 1 & 0 & 0 & 0 & -x'_1x_1 & -x'_1y_1 & -x_1 \\ \vdots & \vdots & \vdots & & & \vdots & \vdots & \vdots \\ x_n & y_n & 1 & 0 & 0 & 0 & -x'_nx_n & -x'_ny_n & -x_n \\ 0 & 0 & 0 & x_1 & y_1 & 1 & -y'_1x_1 & -y'_1y_1 & -y_1 \\ \vdots & \vdots & \vdots & \vdots & & \vdots & \vdots \\ 0 & 0 & 0 & x_n & y_n & 1 & -y'_nx_n & -y'_ny_n & -y_n \end{bmatrix}$$

This problem can be easily solved using Lagrange multipliers:

p is the eigenvector of matrix M<sup>T</sup>M associated to the smallest eigenvalue.

The whole algorithm can be written in 1 (long) line of Matlab!

### proof

#### Lagrangian function

$$L = E' - \lambda(p^T p - 1) = p^T M^T M p - \lambda(p^T p - 1)$$

$$\frac{dL}{dp} = 0 \implies M^T M p - \lambda p = 0$$

$$M^T Mp = \lambda p$$

#### p is na eigen vector of matrix M<sup>T</sup>M

which one? 
$$E = p^T M^T M p = \lambda p^T p = \lambda$$

choose  $\lambda_{min}$ 

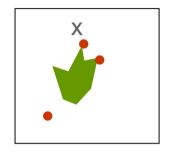
### other transformations?

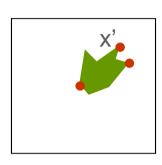
The other transformations (translation, affine, polynomial) are easily estimated by the minimization of the SSD criterion E.

Only the rigid body transformation is a bit more difficult because matrix R is not free. It is a rotation matrix: R<sup>T</sup>R=RR<sup>T</sup>=I and the SSD criterion must be opyimized under this restriction.

This problem can be solved using the singular vector decomposition of the data.

### unknown correspondence





This is a difficult problem!

We need to estimate a permutation matrix.



$$p = \begin{bmatrix} 0 & 0 & 0 \\ 1 & 0 & 0 \\ 0 & 1 & 0 \end{bmatrix}$$

which minimizes the matching criterion E.

tough!

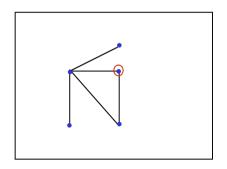
See the paper by Maciel & Costeira, PAMI03

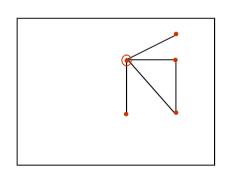
suboptimal approaches are used instead!

### ransac

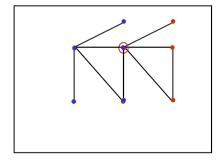
RANSAC stands for Random Sample Consensus (Fischler, Bolles, 1981)

It is based on hypothesis generation and classification of data points as inliers and outliers.





estimate translation



only 2 points are matched! bad attempt!

## ransac (2)

Objective: to estimate a transform W(x,q) with 2n degrees of freedom.

#### **Algorithm**

#### Hypotheses generation

randomly select n pairs of points  $(x_i, x'_k)$ 

estimate the geometric transformation  $W(x,\theta)$ 

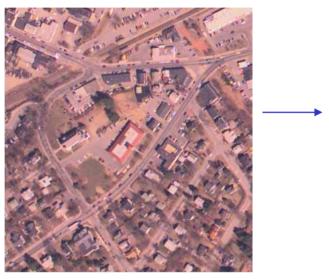
Compute the number of points which were correctly aligned (support) i.e., such that

$$|\mathbf{x'}_k - \mathbf{W}(\mathbf{x}_i, \mathbf{\theta})| < \varepsilon$$

Model selection: choose the transformation with largest support

Refinement: improve the estimate of  $\theta$  by applying the least squares method to the subset of points which are well aligned.

# example - registration





#### Afine transform

(3 marks)

(Matlab demo)



Jorge Marques, 2008

# example - mosaicing





### homography

(4 marks)

# exemplo (cont.)



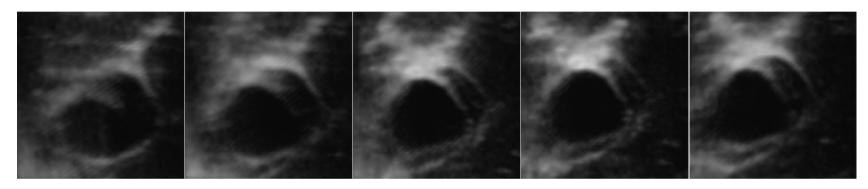
# mosaicing



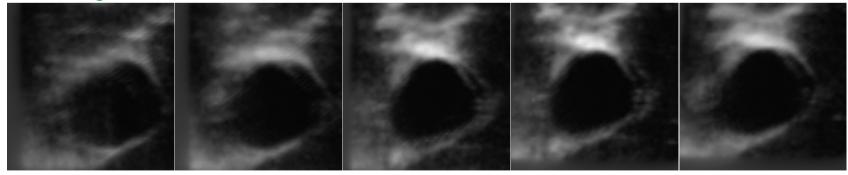
mosaicing → alignment + fusion

## 3D ultrasound

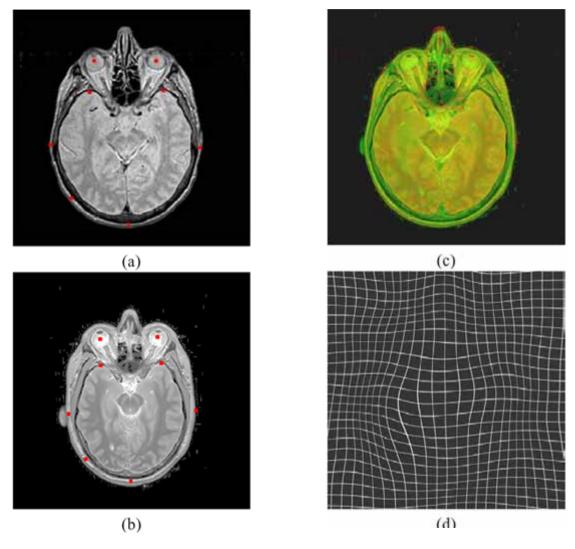
### without alignment



### with alignment



# non-rigid alignment





region tracking



Two steps

region detection region tracking

## Region detection

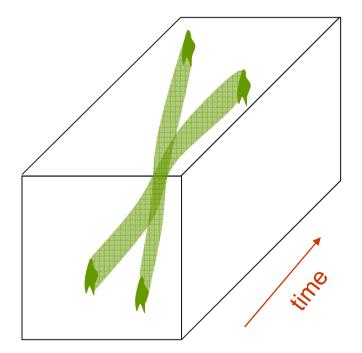
# problem

#### goal:

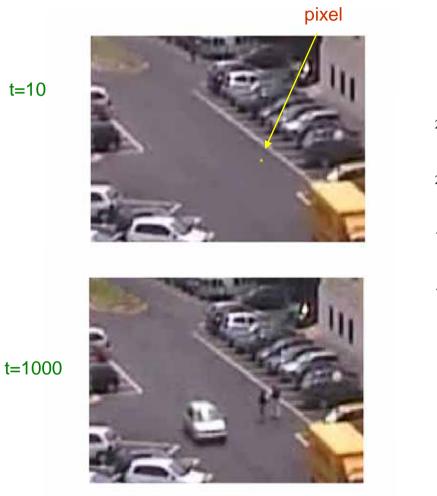
• detect all moving objects

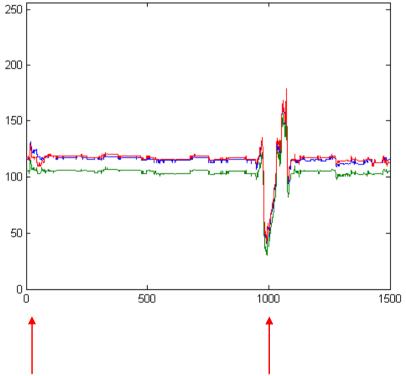
#### assumptions

- static camera
- static background
- show illumination changes



# Evolution of pixel color



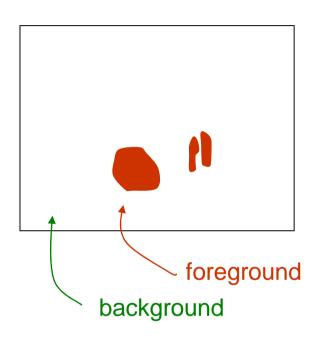


# **Background subtraction**





background image



#### **Pixel classification**

If  $|I(x,y)-B(x,y)| < \epsilon$ , the pixel is classified as background pixel. Otherwise it is classified as active.

### Basic background subtraction

The basic background subtraction classifies a pixel I(x,y) as active if

$$A(x, y) = 1$$
 if  $|I(x, y) - B(x, y)| > \lambda$   
 $A(x, y) = 0$  otherwise

Image A(x,y) is very noisy. It has many small regions classified as active and some true objects appear fragmented in several regions.

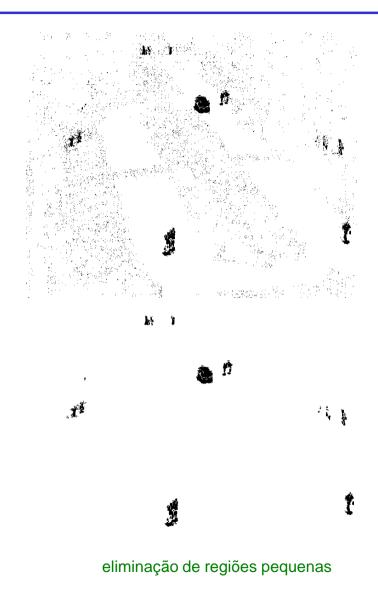
Morphologinal post-processing is usually done. Typically we compute all conected components and eliminate all the small regions.

# Example







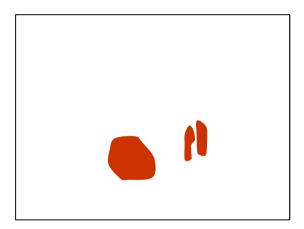


Jorge Marques, 2008

## How to deal with time-varying illumination?

Illumination changes can be compensated by the adaptation of the background image.

Only the pixels belonging to the background regiion should be adapted.



$$B(x, y, t) = \alpha B(x, y, t - 1) + (1 - \alpha)I(x, y, t)$$

background pixels

$$B(x, y, t) = B(x, y, t - 1)$$

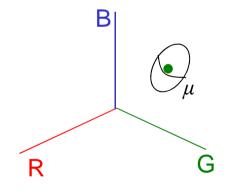
foreground pixels

### Gaussian background model

(see Wren et al., 1997)

Cackground pixels are corrupted by noise. We can model each pixel as a random variable with Gaussian distribution

$$I(x, y) \sim N(\mu(x, y), R(x, y))$$



pixel classification

$$p(I(x, y)) \ge \lambda$$
  $\Rightarrow$  background pixel

$$p(I(x, y)) < \lambda$$
  $\Rightarrow$  foreground pixel

$$p(I(x,y)) = \frac{1}{(2\pi)^{3/2} \det(R)^{1/2}} e^{-\frac{1}{2}(I(x,y) - \mu(x,y))^T R^{-1}(I(x,y) - \mu(x,y))}$$

#### Estimation of the Gaussian model

#### batch

$$\mu(x,y) = \frac{1}{T} \sum_{t=1}^{T} I(x,y,t)$$

$$R(x,y) = \frac{1}{T} \sum_{t=1}^{T} (I(x,y,t) - \mu(x,y)) (I(x,y,t) - \mu(x,y))^{T}$$

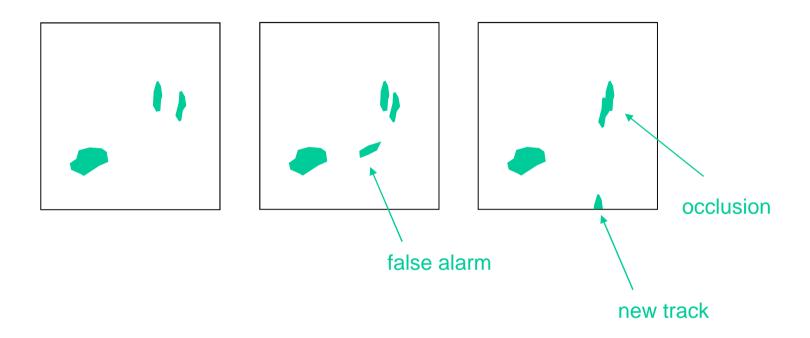
#### adaptive

$$\mu(x, y, t) = \alpha \mu(x, y, t - 1) + (1 - \alpha)I(x, y, t)$$

$$R(x, y, t) = \alpha R(x, y, t - 1) + (1 - \alpha)(I(x, y, t) - \mu(x, y, t - 1))(I(x, y, t) - \mu(x, y, t - 1))^{T}$$

### region tracking

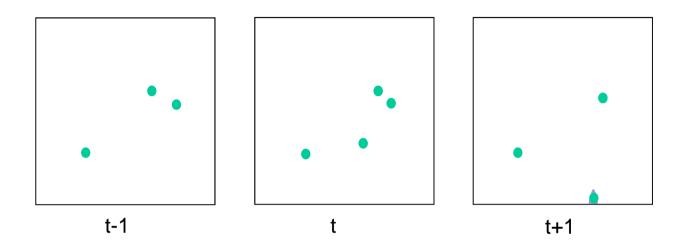
### region tracking



Goal: find the trajectory of each object along multiple frames

Dificulties: misdetections, false alarms, occlusions, object splits and merges, new tracks

## point tracking



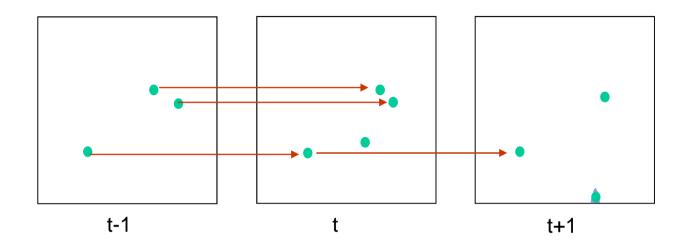
**Data**  $D = \{(t, p_i^t)\}$ 

 $p_i^t$  position of the i-th region at frame t

Track is a sequence of points detected at different (usually consecutive) frames

$$T = \{(t_1, x_1), (t_2, x_2), (t_n, x_n)\} \qquad (t_i, x_i) \in D, \quad t_i < t_{i+1}$$
 
$$(t_{i+1} = t_i + 1)$$

#### point association



#### available methods:

Statistical: propagate uncertainty and assume a dynamic model for the target trajectories (e.g., Kalman or PDA filter)

Deterministic: based on assignment costs and do not require dynamic models (e.g., graph based methods)

### hypotheses

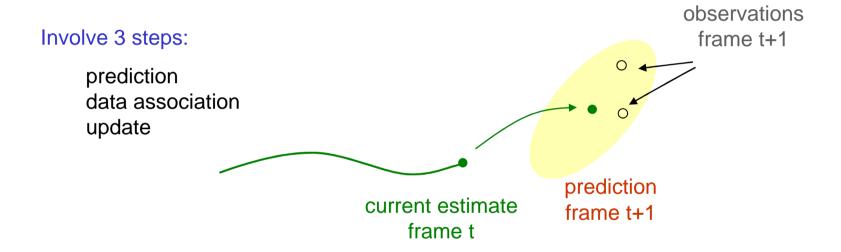


#### typical assumptions

- (a) only regions detected in consecutive frames can be associated
- (b) regions should correspond to a single target (and vice-versa)
- (c) new objects may appear (track birth)
- (d) objects can disapear or be occluded (track death)
- (b') objects can overlap and form groups

#### statistical methods

Statistical methods assume we know a set of tracks and wish to extend them in new frames.



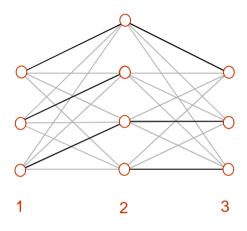
#### Difficulties:

data association problem initialization of new tracks

#### Methods:

nearest–neighbor Kalman filter probabilistic data association filter joint probabilistic data association filter particle filter

#### methods based on graphs

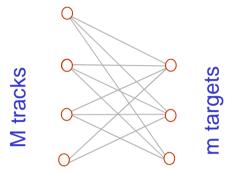


Nodes coorespond to the detected objects in each frame and the links define a solution for the association problem

Each admissible link has a cost C<sub>t</sub>(i,j) (unconnected nodes also have a cost).

#### Veenman et al

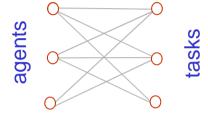
(PAMI 2001)



This method deals with pairs of frames and formulates the association of targets to existing tracks as an assignment problem if M=m.

#### assignment problem

Problem: there are M agents and m tasks (M=m); we wish to assign one agent to one task minimizing the total cost



$$C = \sum_{i,i=1}^{m} a_{ij} c_{ij}$$

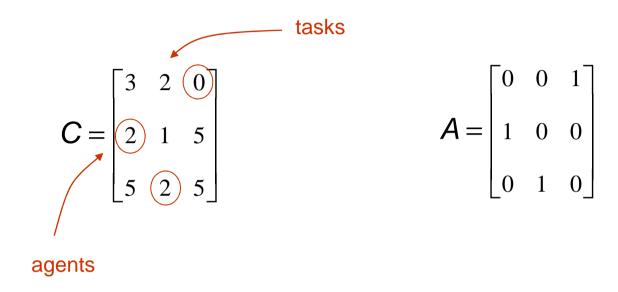
Restrictions

$$\sum_{i=1}^{m} a_{ij} = \sum_{j=1}^{m} a_{ij} = 1 \qquad a_{ij} \in \{0,1\}$$

 $c_{ij}$  is the cost of assigning agent i to task j and  $a_{ij}$  is a binary variable wich is equal to 1 if and only if agent i is assigned to task j.

The minimization of C under these restrictions is a linear programming problem for which there are very efficient algorithms e.g., Hungarian method.

# Example



total cost: 0+2+2=4

#### cost matrix

In tracking, the association cost can be defined in different ways. Two popular choices are

$$a_{ij} = \mid\mid p_i^{t-1} - p_j^t \mid\mid$$

$$a_{ij} = || p_i^{t-1} + v_i^{t-1} - p_j^t ||$$

 $v_i^{t-1}$  displacement vector computed from a previous assignment. (cannot be used in track initialization)

#### Birth and death of tracks

The previous method does not account for new tracks but it has been extended to allow birth and death of tracks

Consider a problem in which all the targets are new. In this case, all the M tracks should die are all the m targets correspond to new tracks.

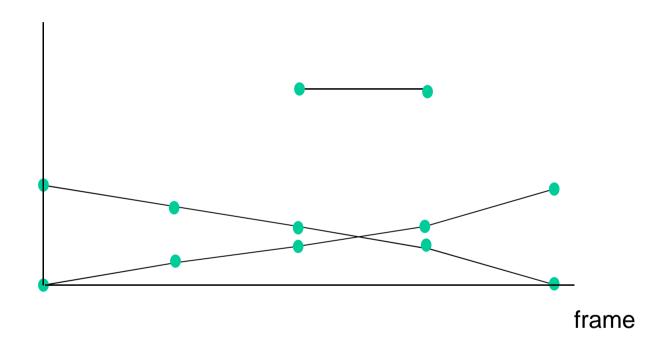
How can we do this in the previous framework?

solution: add M virtual targets and m virtual tracks

 $\mathsf{M} \qquad \mathsf{C} \qquad \qquad c_{ij} = c$ 

$$c_{ij} = c_{high}$$
 if  $i > M$  or  $j > m$ 

# Example 1d



costs were computed using the prediction error, except at the beginning of each track.